

FIG. 1A ~~A~~

PAGE MDLEKTRSERGDGSDVKEKTPPNPKHAKTKEAGDGQP  
 : | | | | : | : | | | | | : |  
 GAGE2 QGHPQTGCECEDGPDGQEMDPPNPNPEEVKTPEEGEKQSQC  
 GAGE1 M MR HYVA..  
 GAGE3-5 M  
 GAGE6 V  
 : | | : : | | : | : |  
 MAGE5 RQSIKSSNQEEEGPSTSPDPE...ESVFRAALSKKKVADLIHFLLLKY  
 MAGE8 S DE SN AHL L E DE E VR R...

FIG. 1B

~~B~~

PAGE1 DLEKTRSERGDGSDVKEKTPPNPKHARTKEAGDQOP  
:  
PAGE2 ALLKIEDEPGDGPDVREGIMPTFDLTKVLEAGDAQP  
:  
PAGE3 TRPKTGGERGDPNVKGESLPNLEFPVKIPEAGEGQPSV

Annotated Sheets

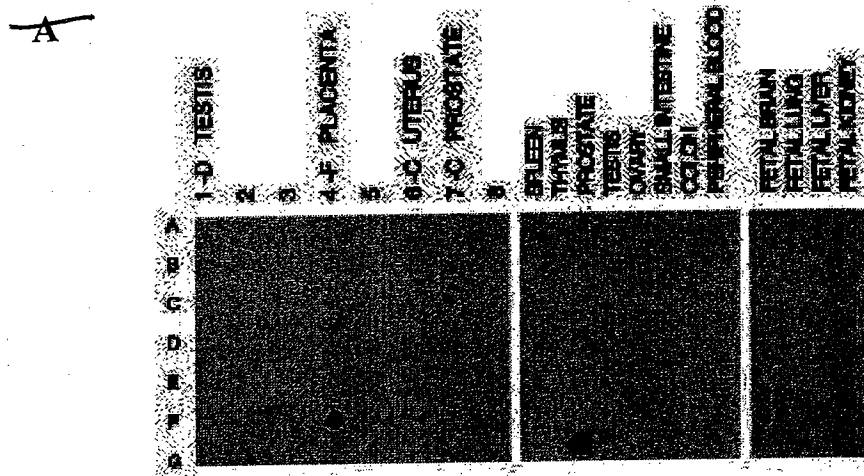


FIG. 2A

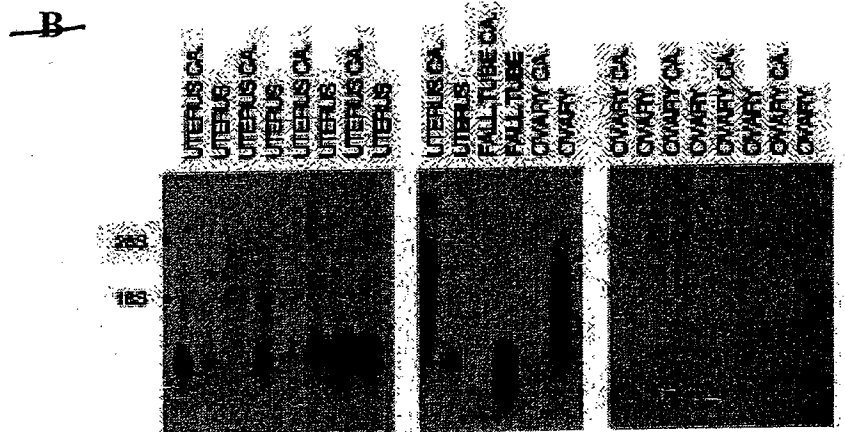
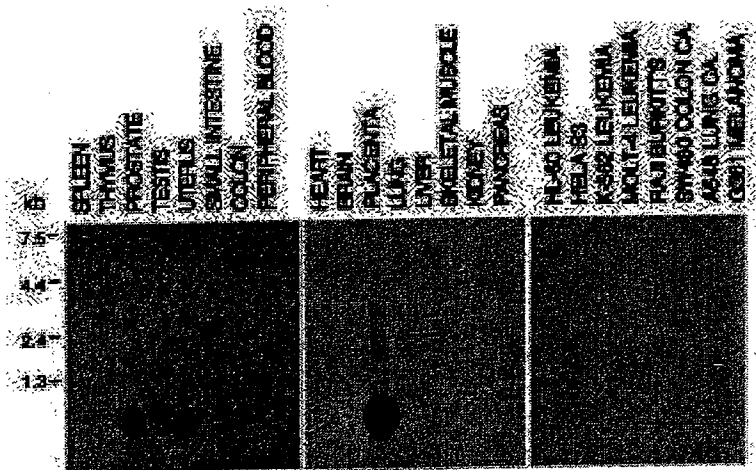


FIG. 2B

Annotated Sheets

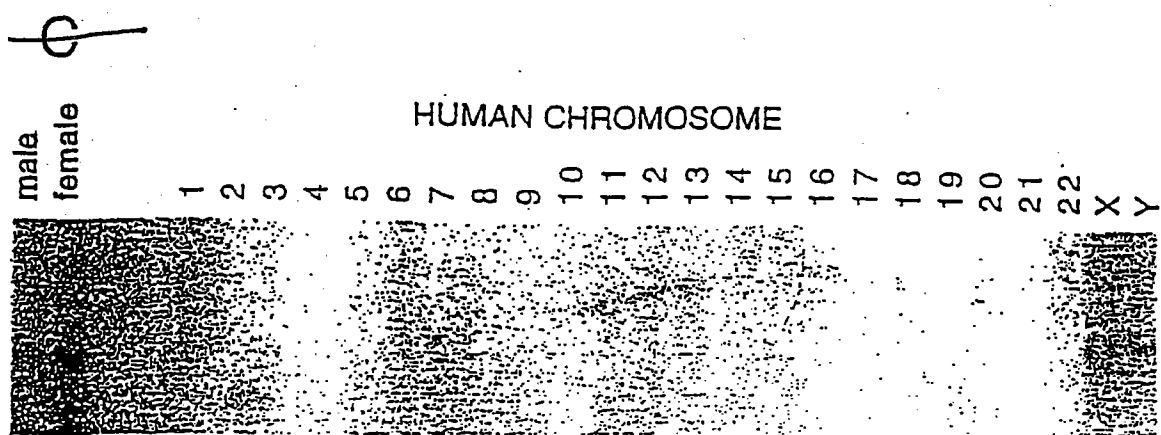


FIG. 2C

~~FIG. 3~~

Annotated Sheets

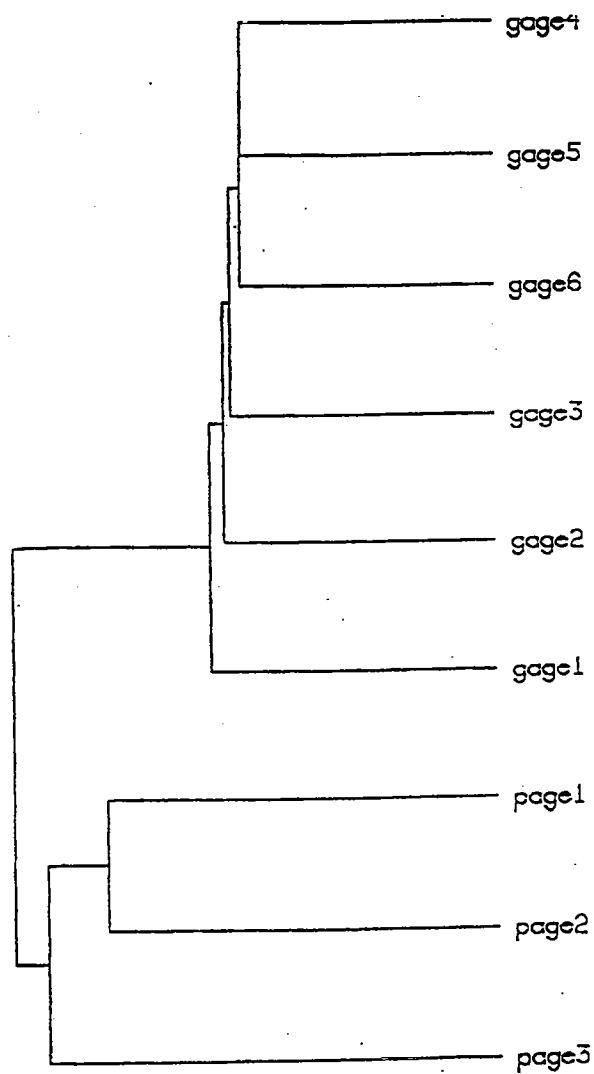


FIG. 3

~~FIG. 4~~

Annotated Sheets

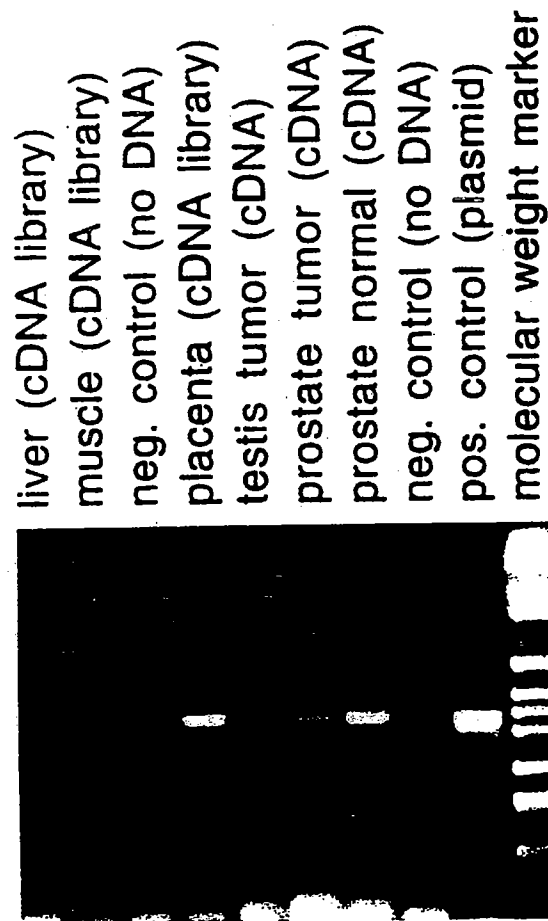


FIG. 4

~~FIG. 5~~

Annotated Sheets

Page 4 sequence (bold and underline: open reading frame).

GAAGAATTCGCCAGGCTCTCTGCTGACTCAAGTTCTTCAGTTCACGATCTTCTAGTT  
GCAGCG

ATGAGTGCACGAGTGAGATCAAGATCCAGAGGAAGAGGAGATGGTCAGGAGGCTCCC  
GATGTGGTTGCATTCGTGGCTCCCGGTGAATCTCAGCAAGAGGAACCACCAACTGAC  
AATCAGGATATTGAACCTGGACAAGAGAGAGAAGGAACACCTCCGATCGAAGAACGT  
AAAGTAGAAGGTGATTGCCAGGAAATGGATCTGGAAAAGACTCGGAGTGAGCGTGGA  
GATGGCTCTGATGTAAAAGAGAAGACTCCACCTAATCCTAAGCATGCTAAGACTAAA  
GAAGCAGGAGATGGGCAGCCA

TAAGTTAAAAGAAGACAAGCTGAAGCTACACACATGGCTGATGTCACATT  
GAAAATGTGACTGAAAATTTGAAAATTTCTCTCAATAAAGTTTGTAGTTTTCTCTGAA

FIG. 5

FIG. 6